## Amendment To The Claims

The following listing of claims will replace all prior versions, and listings, of claims in the application.

- (Currently amended) A method for generating a genetically modified 1. organism for drug screening, which comprises the steps of:
- causing heterologous expression of at least one protein or protein fragment by a) genetic modification of the organism
- analyzing the modified gene expression pattern and identifying b) compensatingly compensating differentially regulated genes
- phenotyping the organism-wherein phenotyping is carried out by the C) reduction or elimination of compensating differential expression or by the labeling of at least one compensating differentially regulated gene.
  - 2. (Canceled)
- (Currently amended) The method as claimed in either of claims of claim 1 and 2, wherein the genetic modification causes heterologous expression of at least one protein or protein fragment which is endogenous or foreign to the organism and/or foreign.
- (Currently amended) The method as claimed in any of claims of claim 1-to-3, 4. wherein the genetic modification causes reduction or elimination of the expression of at least one protein endogenous to the organism.
- (Currently amended) The method as claimed in any of claims of claim 1 to 4, 5. wherein the modified expression is inducible.
- (Currently amended) The method as claimed in of claim 5, wherein the 6. genetic modification comprises introducing a vector which enables the protein or protein fragment to be inducibly expressed, preferably a vector inducible with galactose, copper tetracycline or other comparably inducible vectors.

- (Currently amended) The method as claimed in any of claims 1 to of claim 6, 7. wherein the genetic modification comprises a knock out, preferably an inducible knock out.
- 8. (Currently amended) The method as plaimed in any of claims 1 to 7 claim 1. wherein the organism is drosophila, C. elegans, a prokaryotic or a eukaryotic cell.
- 9. (Currently amended) The method as claimed in of claim 8, wherein the cell is a yeast cell, preferably a yeast cell of the strain S. cerevisiae.
- (Currently amended) The method as claimed in any of claims 1 to of claim 9, 10. wherein the modified gene expression is analyzed with the aid of DNA or protein microarrays.
- (Currently amended) The method as claimed in any of claims 1 to of claim 11. 10, wherein phenotyping is carried out by reducing or eliminating expression of the compensatingly compensating differentially regulated gene.
- 12. (Currently amended) The method as claimed in of claim 11, wherein expression of the compensatingly compensating differentially expressed gene is enhanced te in control organisms and the reduction or elimination is caused by at least partial inhibition of said enhanced expression.
- (Currently amended) The method as claimed in of claim 7, wherein the knock 13. out of the differentially expressed gene is carried out by {lacuna} replacing at least part of the coding sequence of the differentially regulated gene with the coding sequence of a reporter gene or parts of the reporter gene seguence which are sufficient to be detected.
- (Currently amended) The method as claimed in of claim 11, wherein the differentially expressed gene is less strongly expressed than in control organisms and the reduction or elimination of the differential expression is carried out by enhancing its expression of the differentially expressed gene.
- (Currently amended) The method as claimed in any of claims 1 to of claim 15. 14, wherein the reduction or elimination leads to growth inhibition of the organism.

- 16. (Currently amended) The method as claimed in any of claims 1 to of claim 10, wherein phenotyping is carried out by labeling the gene product of the compensatingly compensating differentially regulated gene.
- 17. (Currently amended) A genetically modified, phonotype phenotyped organism, obtained by a the method of claim 1 as claimed in any of claims 1 to 16.
  - 18. (Currently amended) A genetically modified organism, having
- a) genetically modified expression of at least one endogenous or foreign gene, which results in compensatingly compensating differential expression of at least one other gene endogenous to said organism, and
- b) a phenotype caused by reducing/eliminating the reduction or elimination of the compensatingly compensating differential expression of the gene or by labeling the compensatingly compensating differentially regulated gene product.
  - 19. (Canceled)
- 20. (Currently amended) A method for identifying substances a substance having an effect on the function of the <u>a</u> heterologously expressed protein or protein fragment, which method comprises the <u>steps of:</u>
  - a) contacting said substance with said genetically modified organism of claim

    17 or 18 and
  - b) measuring the change in said modified organism as compared to genetically unmodified organism use of the organism as claimed in either of claims 17 or 18.
- 21. (Currently amended) An assay for drug screening using at least one phenotype phenotyped organism as claimed in either of claims 17 or 18, which comprises the steps of:
  - e a) determining the phenotype of said organism
  - d b) contacting the substance to be tested with said organism

- e <u>c</u>) observing a possible modification of said phenotype.
- (Canceled) 22.